

Pharmacogenetic Activities in SWOG Breast Cancer



Pharmacogenomics: Future Plans

- **S8897 Adjuvant CMF vs. CAF/ no Treatment**
 - Ambrosone **RO1**: Other genes (TBCI approved, analyses ongoing)
- **S0221 Adjuvant Dose Dense vs. Dose Denser AC-T**
 - Ambrosone: These and other genes (TBCI approved, collection ongoing)
 - Toxicities (short run)
 - Outcomes (long run)
- **S0226 Metastatic anastrozole +/- fulvestrant**
 - Susan Nowell **RO1**
 - ESR 1 and 2 (ER)
 - CYP19 (Aromatase)
 - CYP3A4, FMO3, UGT2B3, UGT2B10, SULT1E1 and SULT1A1
- **ALL: Whole genome sequencing (PGRN collaborations)**
(Christine Ambrosone is applying for **RO1** for S0221)

Pharmacogenomics: Future Plans

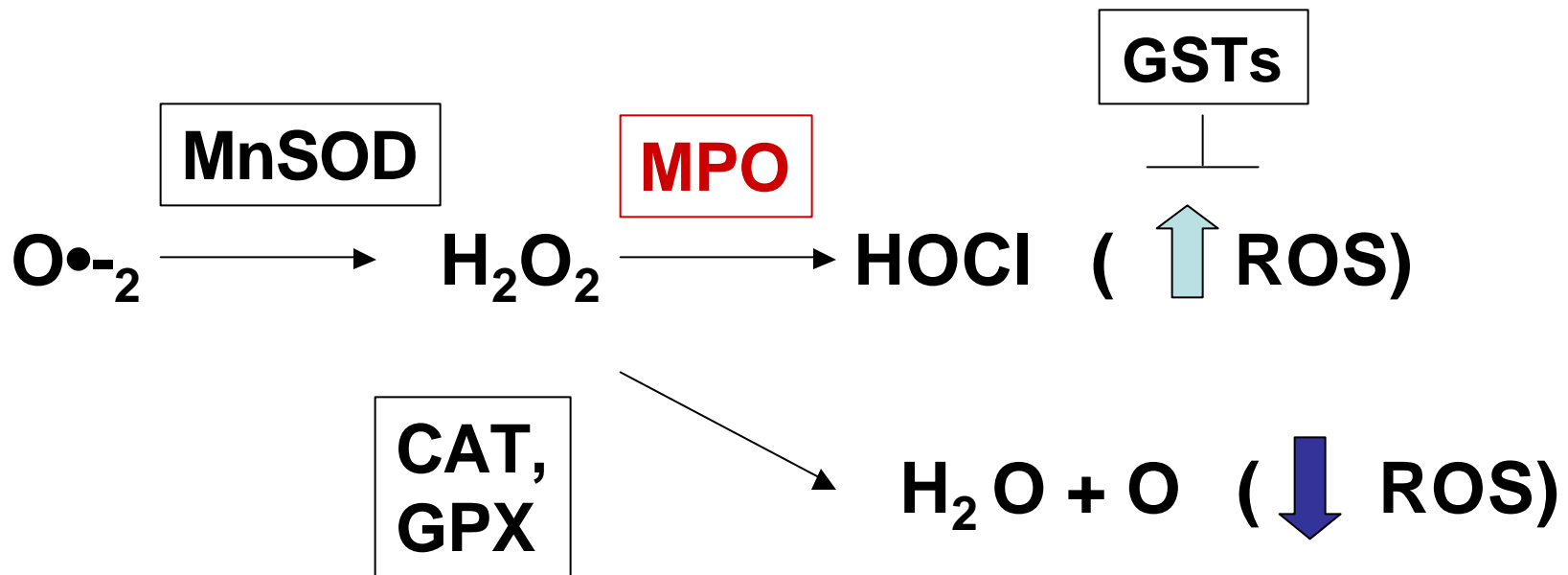
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Inherited genetic variability

Chemotherapy, Radiation and ROS

- **ROS/Oxidative stress induced by CTX:**
 - **Damage to DNA**
 - **Lipid peroxidation**
 - **Protein modification**
 - **Membrane disruption**
 - **Mitochondrial damage**
 - **Apoptotic cascade**
- **Inherited variability in generation of ROS could result in differential treatment outcomes**

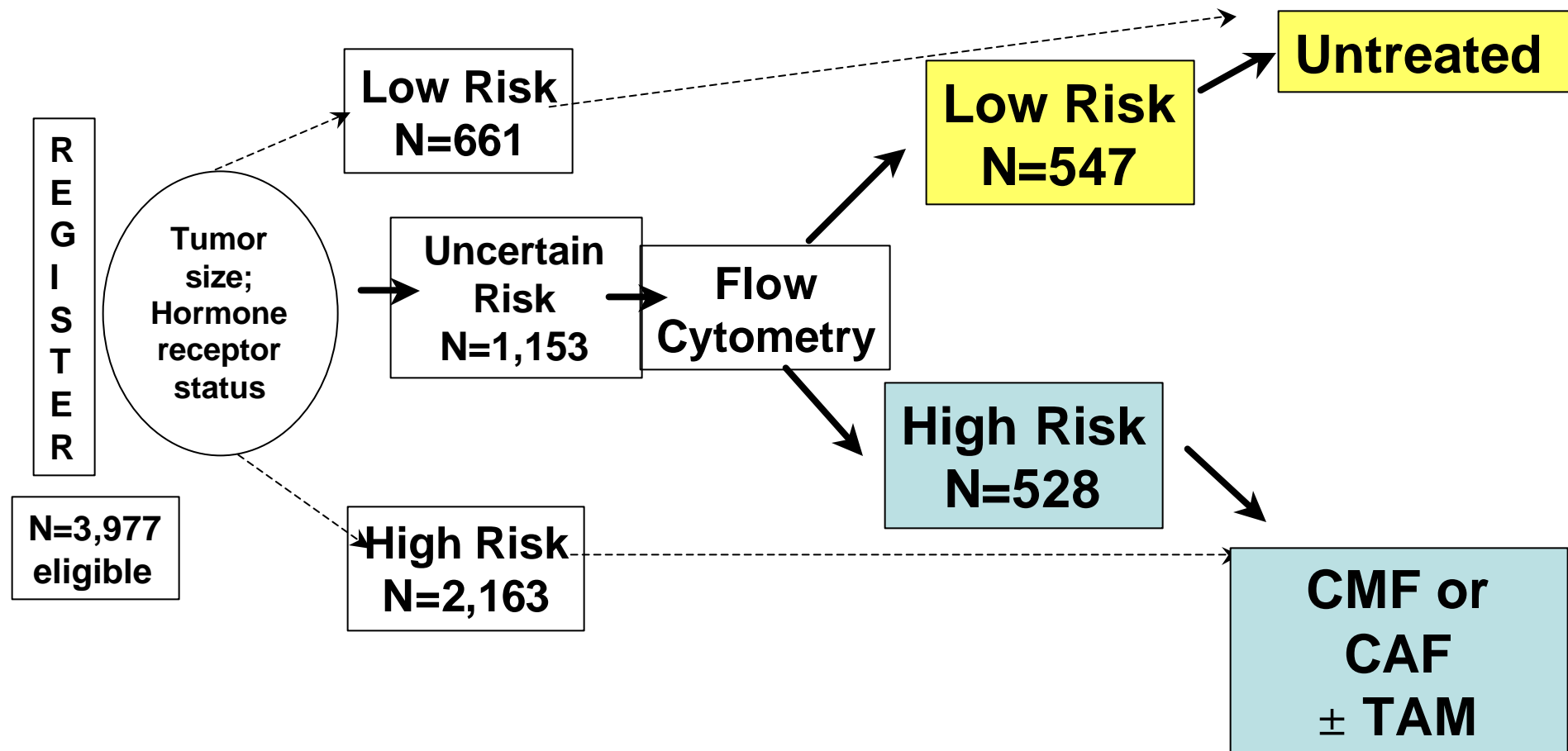
Endogenous Oxidant and Antioxidant Capabilities



- Genes for these enzymes are polymorphic
- SNPs dictate different levels of enzyme activity
- Different genotypes may be sensitive or resistant to chemotherapy

Schema in SWOG S8897

High risk, node negative women (tumor > 2cm or negative ER or high S phase fraction)

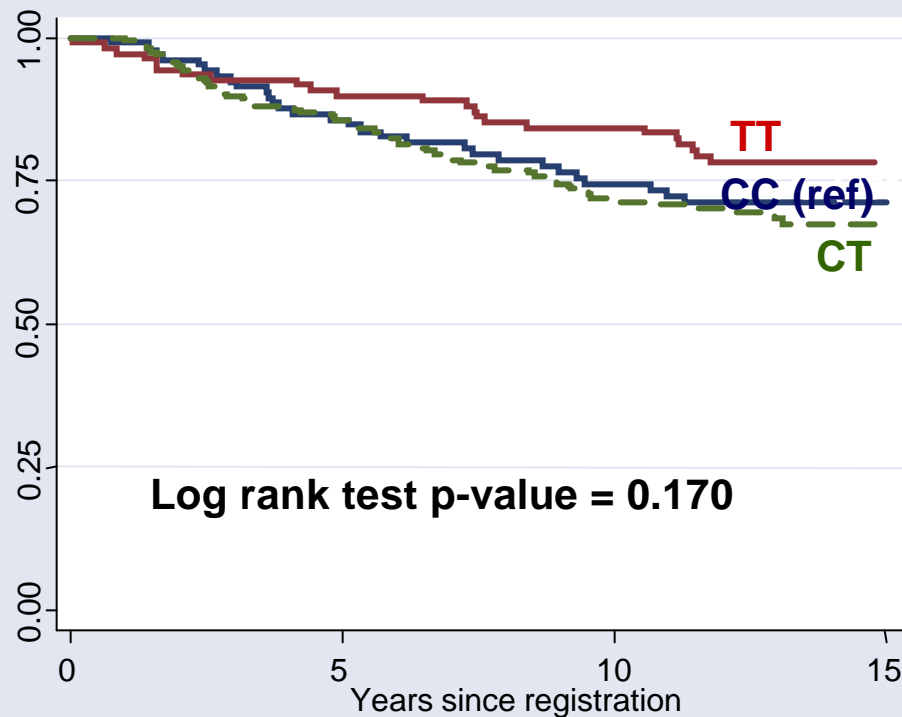


Methods

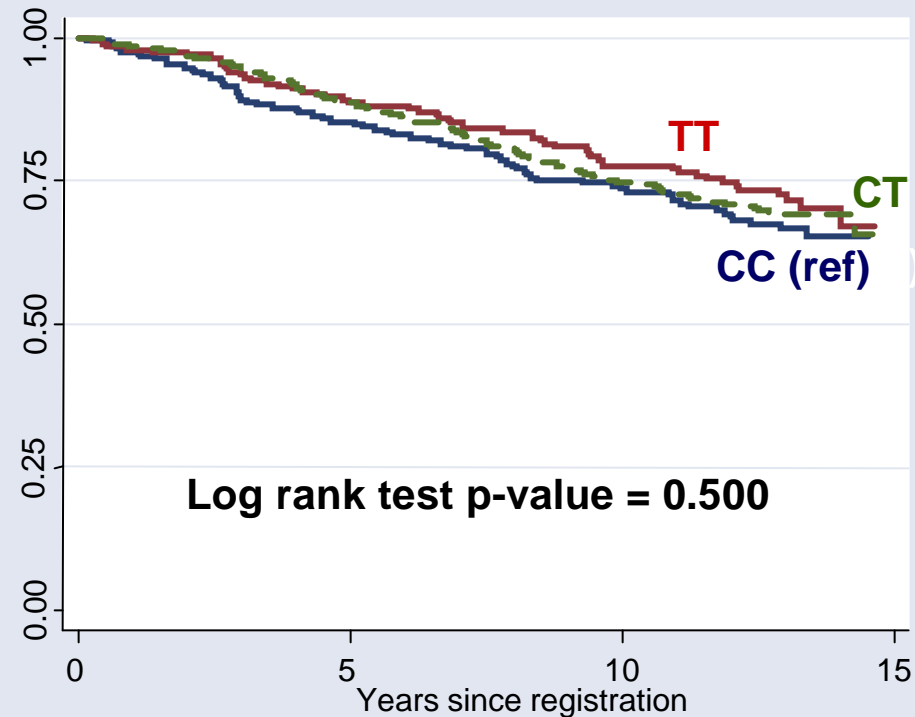
- **Normal lymph node tissue available for genotyping**
 - **Low risk: no treatment**
 - **Intermediate risk: all received chemotherapy (CMF vs. CAF)**
- **Genotyping performed by Ambrosone (RPCI) and Rae laboratories (UMCCC)**

MnSOD C/T (Ala-9Val) and Breast Cancer Disease-Free Survival

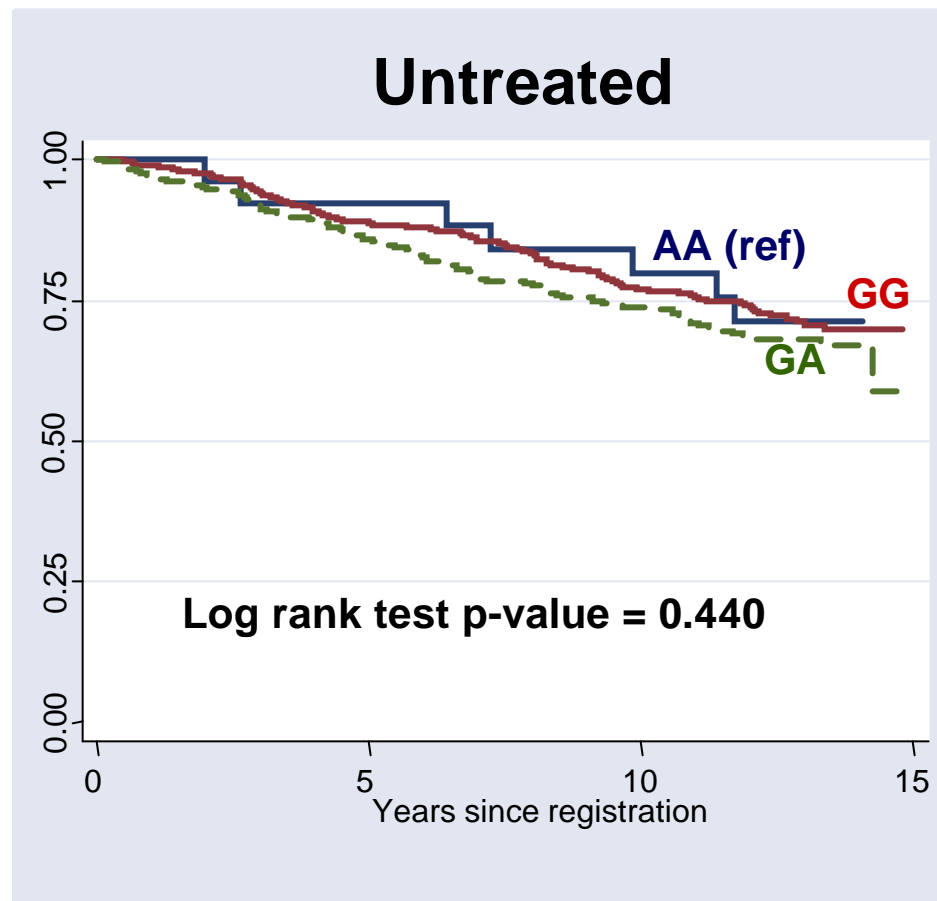
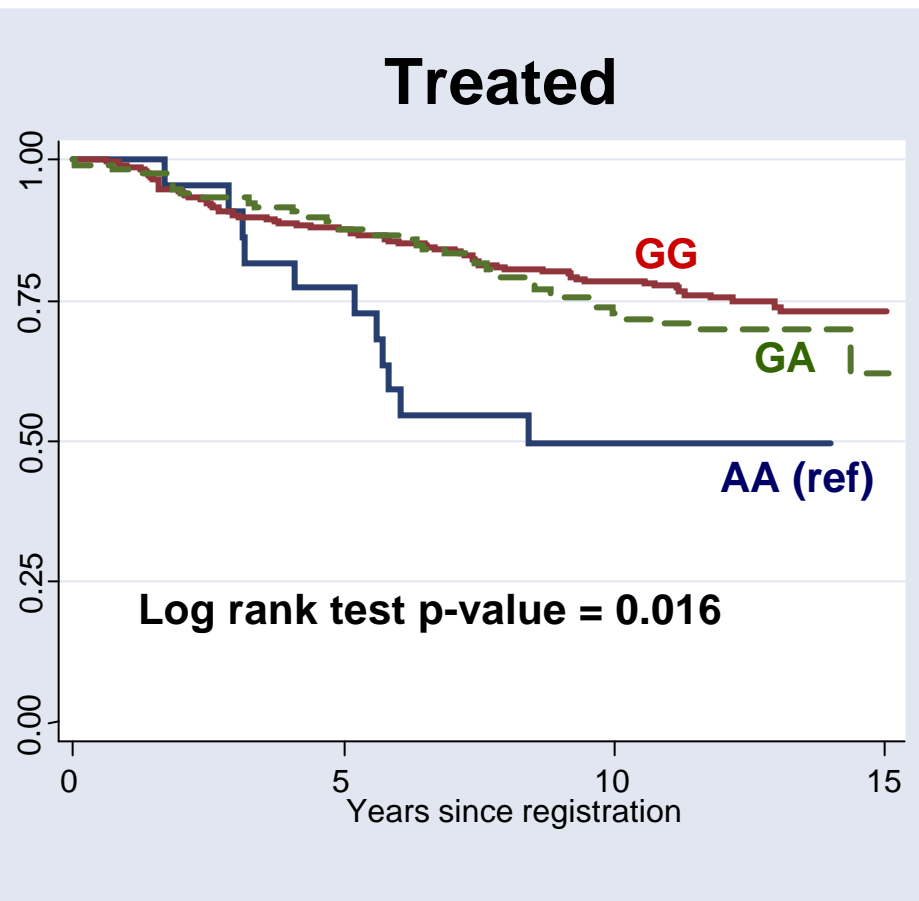
Treated



Untreated



MPO -463 G/A and Breast Cancer Disease-Free Survival



Associations between *MPO* Genotype and DFS, Treated and Untreated Arms

genotype	Treated			Untreated		
	censored	failures	HR (95% CI)	censored	failures	HR (95% CI)
<i>MPO</i>						
AA	11	11	1.0 (ref)	19	7	1.0 (ref)
AG	84	36	0.51 (0.26-0.99)	169	77	1.27 (0.59-2.76)
GG	194	65	0.41 (0.21-0.77)	350	131	1.08 (0.50-2.31)
AA	11	11	1.0 (ref)	19	7	1.0 (ref)
AG+GG	278	101	0.44 (0.23-0.82)	519	208	1.14 (0.54-2.43)

Ambrosone et al San Antonio Breast Cancer Symposium, 2006

HR adjusted for menopausal status, and time between surgery and chemotherapy

Summary and Conclusions

- **High activity *MPO* genotypes are associated with better outcome in women treated with chemotherapy**
 - **Results similar in CMF and CAF groups**
 - **Not PROGNOSTIC (no effect in *untreated* patients)**
- **No effect detected for MnSOD; treated or not**

Limitations

- **Very limited quantities of DNA – only single SNPs assessed**
- **Limited number of genes in pathways examined**
- **Limited sample size**
- **Need for high quality DNA (WBCs) from large clinical trials, comprehensive assessment of variability across genes in pathways**

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**S0221: Phase III Trial of Continuous
Schedule AC+G vs Q2 Week Schedule
AC, Followed by Paclitaxel Given Either
Every 2 Weeks or Weekly for 12 Weeks
as Post-Operative Adjuvant Therapy in
Node-Positive or High-Risk Node
Negative Breast Cancer (Budd, PI)**

Accrual goal – 3,250

- **Collection of 2 tubes of blood from consenting patients (1 red-top banked for banked serum, 1 purple top for DNA extraction)**
- **As of 3/27/08, samples received from 903 patients**
- **Also collection of questionnaire data**
- **Application to TBCI for use of DNA for Genome Wide Study in relation to toxicity and DFS**

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SWOG Breast Cancer Pharmacogenomics

- S0702
 - A prospective registry of cancer patients with metastatic bone disease taking bisphosphonate therapy
 - Endpoint: Osteonecrosis of the Jaw (ONJ)
 - PI: Cathy Van Poznak, Julie Gralow
 - n>7000
 - WBC collected and stored on all

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